

Resource Summary Report

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DISULFIND

RRID:SCR_016072

Type: Tool

Proper Citation

DISULFIND (RRID:SCR_016072)

Resource Information

URL: <http://disulfind.dsi.unifi.it/>

Proper Citation: DISULFIND (RRID:SCR_016072)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on February 28,2023, Software for predicting the disulfide bonding state of cysteines and their disulfide connectivity, starting from a protein sequence alone and may be useful in other genomic annotation tasks.

Abbreviations: Disulfinder

Synonyms: Cysteines Disulfide Bonding State and Connectivity Predictor

Resource Type: data analysis software, software application, software resource, sequence analysis software, data processing software

Defining Citation: [PMID:16844986](#), [DOI:10.1093/nar/gkl266](#)

Keywords: predict, disulfide, bonding, state, cysteine, protein, sequence, genomic, annotation, bio.tools

Funding: EU STREP APriL II contract no. FP6-508861;
EU NoE BIOPATTERN contract no. FP6-508803;
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Engineering and Technology

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: DISULFIND

Resource ID: SCR_016072

Alternate IDs: OMICS_04214, biotools:disulfind

Alternate URLs: <https://bio.tools/disulfind>, <https://sources.debian.org/src/disulfinder/>

Record Creation Time: 20220129T080328+0000

Record Last Update: 20260801T190527+0000

Ratings and Alerts

No rating or validation information has been found for DISULFIND.

No alerts have been found for DISULFIND.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 66 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Salabi F, et al. (2023) Identification, classification, and characterization of alpha and beta subunits of LVP1 protein from the venom gland of four Iranian scorpion species. Scientific reports, 13(1), 22277.

Brewer MS, et al. (2023) Killer Knots: Molecular Evolution of Inhibitor Cystine Knot Toxins in Wandering Spiders (Araneae: Ctenidae). Toxins, 15(2).

Nuryana I, et al. (2022) Expression of Codon-Optimized Gene Encoding Murine Moloney Leukemia Virus Reverse Transcriptase in Escherichia coli. The protein journal, 41(4-5), 515.

Pimkova K, et al. (2022) Quantitative analysis of redox proteome reveals oxidation-sensitive protein thiols acting in fundamental processes of developmental hematopoiesis. Redox biology, 53, 102343.

Arenas NE, et al. (2022) Design of a specific peptide against phenolic glycolipid-1 from Mycobacterium leprae and its implications in leprosy bacilli entry. Memorias do Instituto Oswaldo Cruz, 117, e220025.

Sacchi S, et al. (2021) Yin and Yang in Post-Translational Modifications of Human D-Amino Acid Oxidase. Frontiers in molecular biosciences, 8, 684934.

Estrada-Gómez S, et al. (2021) Analysis of High Molecular Mass Compounds from the Spider Pamphobeteus verdolaga Venom Gland. A Transcriptomic and MS ID Approach. Toxins, 13(7).

Zhou X, et al. (2021) Expression and Function Analysis of Interleukin-17A/F1, 2, and 3 Genes in Yellow Catfish (Pelteobagrus fulvidraco): Distinct Bioactivity of Recombinant IL-

17A/F1, 2, and 3. *Frontiers in immunology*, 12, 626895.

Sadri Najafabadi Z, et al. (2021) Designing of a chimeric protein contains StxB, intimin and EscC against toxicity and adherence of enterohemorrhagic *Escherichia coli* O157:H7 and evaluation of serum antibody titers against it. *Molecular immunology*, 134, 218.

Bartošová-Sojková P, et al. (2021) Evolutionary Analysis of Cystatins of Early-Emerging Metazoans Reveals a Novel Subtype in Parasitic Cnidarians. *Biology*, 10(2).

Ebert MK, et al. (2021) Identification and characterization of *Cercospora beticola* necrosis-inducing effector CbNip1. *Molecular plant pathology*, 22(3), 301.

Holzknecht J, et al. (2020) The *Penicillium chrysogenum* Q176 Antimicrobial Protein PAFC Effectively Inhibits the Growth of the Opportunistic Human Pathogen *Candida albicans*. *Journal of fungi (Basel, Switzerland)*, 6(3).

Chen C, et al. (2020) Multimodal imaging and genetic characteristics of Chinese patients with USH2A-associated nonsyndromic retinitis pigmentosa. *Molecular genetics & genomic medicine*, 8(11), e1479.

Nikoloudakis N, et al. (2020) Structural Diversity and Highly Specific Host-Pathogen Transcriptional Regulation of Defensin Genes Is Revealed in Tomato. *International journal of molecular sciences*, 21(24).

Joo MS, et al. (2020) The molecular characterization, expression analysis and antimicrobial activity of theromacin from Asian polychaeta (*Perinereis lineata*). *Developmental and comparative immunology*, 112, 103773.

Christie AE, et al. (2020) Multiple transcriptome mining coupled with tissue specific molecular cloning and mass spectrometry provide insights into agatoxin-like peptide conservation in decapod crustaceans. *General and comparative endocrinology*, 299, 113609.

Fu T, et al. (2020) Two antimicrobial genes from *Aegilops tauschii* Cosson identified by the *Bacillus subtilis* expression system. *Scientific reports*, 10(1), 13346.

Deng M, et al. (2020) Jingzhaotoxin-X, a gating modifier of Kv4.2 and Kv4.3 potassium channels purified from the venom of the Chinese tarantula *Chilobrachys jingzhao*. *The journal of venomous animals and toxins including tropical diseases*, 26, e20190043.

Su Y, et al. (2020) A novel C-type lectin with a YPD motif from *Portunus trituberculatus* (PtCLec1) mediating pathogen recognition and opsonization. *Developmental and comparative immunology*, 106, 103609.

Cui ZW, et al. (2020) Splicing variants of grass carp (*Ctenopharyngodon idellus*) IL-21: Functions in IgM+ B cell proliferation and IgM secretion. *Developmental and comparative immunology*, 110, 103728.